IN THE CLAIMS:

Please cancel claims 1-18, 20, and 22-24.

- 1. (CANCELED)
- 2. (CANCELED)
- 3. (CANCELED)
- 4. (CANCELED)
- (CANCELED)
- 6. (CANCELED)
- 7. (CANCELED)
- 8. (CANCELED)
- 9. (CANCELED)
- 10. (CANCELED)
- 11. (CANCELED)
- 12. (CANCELED)
- 13. (CANCELED)
- 14. (CANCELED)
- 15. (CANCELED)
- 16. (CANCELED)
- 17. (CANCELED)
- 18. (CANCELED)

FINNEGAN HENDERSON FARABOW GARRETT & DUNNER

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- 19. (AMENDED) A method of identifying agonists to the IGS1 polypeptide efclaim-13 comprising an amino acid sequence that is at least 80% identical to the amino acid sequence of SEQ ID NO: 2 over its entire length, comprising:
- (a) contacting a cell which produces a IGS1 polypeptide with a test compound; and
- (b) determining whether the test compound effects a signal generated by activation of the IGS1 polypeptide.
 - 20. (ORIGINAL) An agonist identified by the method of claim 19.
- 21. (AMENDED) The method for identifying antagonists to the IGS1 polypeptide of claim-13 comprising an amino acid sequence that is at least 80% identical to the amino acid sequence of SEQ ID NO: 2 over its entire length, comprising:
 - (a) contacting a cell which produces a IGS1 polypeptide with an agonist; and
- (b) determining whether the signal generated by said agonist is diminished in the presence of a candidate compound.
 - 22. (ORIGINAL) An antagonist identified by the method of claim 21.
 - 23. (CANCELED)
 - 24. (CANCELED)
- 25. (NEW) The method of claim 19 wherein the IGS1 polypeptide comprises an amino acid sequence identical to the amino acid sequence of SEQ ID NO: 2.
- 26 (NEW) The method of claim 21 wherein the IGS1 polypeptide comprises an amino acid sequence identical to the amino acid sequence of SEQ ID NO: 2.

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Claims

1.	An isolated polynucleotide	comprising	а	nucleotide	sequence	selected	from	the	group
	consisting of:								

- a nucleotide sequence encoding the IGS1 polypeptide according to SEQ ID NO:
 2;
 - a nucleotide sequence encoding the polypeptide encoded by the DNA insert contained in the deposit no. CBS 102049 at the Centraalbureau voor Schimmelcultures at Baam the Netherlands, in particular a nucleotide sequence corresponding to the SEQ ID NO: 1;
 - a nucleotide sequence having at least 80 % (preferably at least 90%) sequence identity over its entire length to the nucleotide sequence of (a) or (b);
 - a nucleotide sequence which is complimentary to the nucleotide sequence of (a) or (b) or (c).
- The polynucleotide of claim 1 wherein said polynucleotide comprises the nucleotide sequence contained in SEQ ID NO:1 encoding the IGS1 polypeptide of SEQ ID NO:2.
- The polynucleotide of claim 1 wherein said polynucleotide comprises a nucleotide
 sequence that is at least 80% identical to that of SEQ ID NO:1 over its entire length.
 - 4. The polynucleotide of claim 3 which is the polynucleotide of SEQ ID NO:1.
 - The polynucleotide of claim 1-4 which is DNA or RNA.
 - 6. A DNA or RNA molecule comprising an expression system, wherein said expression system is capable of producing an IGS1 polypeptide comprising an amino acid sequence, which has at least 80% identity with the polypeptide of SEQ ID NO:2 when said expression system is present in a compatible host cell.
 - A host cell comprising the expression system of claim 6.
 - 8. A host cell according to claim 7 which is a yeast cell
- 35 9. A host cell according to claim 7 which is an animal cell
 - 10. IGS1 receptor membrane preparation derived from a cell according to claim 7-9.

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- 11. A process for producing an IGS1 polypeptide comprising culturing a host of claim 7 under conditions sufficient for the production of said polypeptide and recovering the polypeptide from the culture.
- 5 12. A process for producing a cell which produces an IGS1 polypeptide thereof comprising transforming or transfecting a cell with the expression system of claim 6 such that the cell, under appropriate culture conditions, is capable of producing an IGS1 polypeptide.
- An IGS1 polypeptide comprising an amino acid sequence which is at least 80% identical to
 the amino acid sequence of SEQ ID NO:2 over its entire length.
 - 14. The polypeptide of claim 13 which comprises the amino acid sequence of SEQ ID NO:2.
 - 15. An antibody immunospecific for the IGS1 polypeptide of claim 13.
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 - 16. A method for the treatment of a subject in need of enhanced activity or expression of IGS1 polypeptide receptor of claim 13 comprising:
 - administering to the subject a therapeutically effective amount of an agonist to said receptor; and/or
 - (b) providing to the subject an isolated polynucleotide comprising a nucleotide sequence that has at least 80% identity to a nucleotide sequence encoding the IGS1 polypeptide of SEQ ID NO:2 over its entire length; or a nucleotide sequence complementary to said nucleotide sequence in a form so as to effect production of said receptor activity in vivo.

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- A method for the treatment of a subject having need to inhibit activity or expression of IGS1 polypeptide receptor of claim 13 comprising;
 - administering to the subject a therapeutically effective amount of an antagonist to said receptor; and/or
 - administering to the subject a polynucleotide that inhibits the expression of the nucleotide sequence encoding said receptor; and/or
 - administering to the subject a therapeutically effective amount of a polypeptide that competes with said receptor for its ligand.
- 35 18. A process for diagnosing a disease or a susceptibility to a disease in a subject related to expression or activity of the IGS1 polypeptide of claim 13 in a subject comprising:
 - determining the presence or absence of a mutation in the nucleotide sequence encoding said IGS1 polypeptide in the genome of said subject; and/or

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- (b) analyzing for the presence or amount of the IGS1 polypeptide expression in a sample derived from said subject.
- 19. A method for identifying agonists to the IGS1 polypeptide of claim 13 comprising:
 - (a) contacting a cell which produces a IGS1 polypeptide with a test compound; and
 - (b) determining whether the test compound effects a signal generated by activation of the IGS1 polypeptide.
- An agonist identified by the method of claim 19.

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- 21. The method for identifying antagonists to the IGS1 polypeptide of claim 13 comprising:
 - (a) contacting a cell which produces a IGS1 polypeptide with an agonist; and
 - (b) determining whether the signal generated by said agonist is diminished in the presence of a candidate compound.
- 22. An antagonist identified by the method of claim 21.
- A recombinant host cell produced by a method of claim 12 or a membrane thereof expressing an IGS1 polypeptide.
- 24. A method of creating a genetically modified non-human animal comprising the steps of
 - a) ligating the coding portion of a polynucleotide consisting essentially of a nucleic acid sequence encoding a protein having the amino acid sequence SEQ ID NO: 2 or a biologically active fragment thereof to a regulatory sequence which is capable of driving high level gene expression or expression in a cell type in which the gene is not normally expressed in said animal; or
 - b) engineering the coding portion of a polynucleotide consisting essentially of a nucleic acid sequence encoding a protein having the amino acid sequence SEQ ID NO: 2 or a biologically active fragment thereof and reintroducing said sequence in the genome of said animal in such a way that the endogenous gene alleles encoding a protein having the amino acid sequence SEQ ID NO: 2 or a biologically active fragment are fully or partially inactivated.